

1589.0160001Seq. Listing.txt SEQUENCE LISTING

<110> Wild et al. <120> Methods of Eliciting Broadly Neutralizing Antibodies Targeting HIV-1 gp41 <130> 1589.0160001 <140> <141> <160> 84 <170> PatentIn Ver. 2.1 <210> 1 <211> 55 <212> PRT <213> Human immunodeficiency virus type 1 <400> 1 Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu Thr Val Trp Gly 20 Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu Lys 35 Asp Gln Gln Leu Leu Gly Ile 50 55 <210> 2 <211> 38 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic <400>2Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu 1 5 10 15

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Arg Tyr Leu Lys Asp Gln 35

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20 25 30

Ala Arg Ile Leu 35

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1 5 10 15

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20 25 30

Lys Asn Glu Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp
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Asn Trp Phe Asn Ile Thr Asn Trp 50 55

<210> 5 <211> 36

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Leu Leu
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Page 3

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Ile Val Asn Arg Val Arg Gln Gly Tyr Ser Pro Leu Ser Phe Gln Thr

His Leu Pro Thr Pro Arg Gly Pro Asp Arg Pro Glu Gly Ile Glu Glu

Glu Gly Gly Glu Arg Asp Arg Asp Arg Ser Ile Arg Leu Val Asn Gly

Ser Leu Ala Leu Ile Trp Asp Asp Leu Arg Ser Leu Cys Leu Phe Ser 245 250 255

Tyr His Arg Leu Arg Asp Leu Leu Leu Ile Val Thr Arg Ile Val Glu 260 265 270

Leu Leu Gly Arg Arg Gly Trp Glu Ala Leu Lys Tyr Trp Trp Asn Leu 275 280 285

Leu Gln Tyr Trp Ser Gln Glu Leu Lys Asn Ser Ala Val Ser Leu Leu 290 295 300

Asn Ala Thr Ala Ile Ala Val Ala Glu Gly Thr Asp Arg Val Ile Glu 305 310 315 320

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35 40 45

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Ala Arg Val Leu 35

<210> 15

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Arg Tyr Leu Gly Asp Gln 35

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Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
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Ala Arg Val Leu Ala Leu Glu Arg Tyr Leu Arg Asp Gln
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Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln 20 25 30

Ala Arg Val Leu

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Arg Tyr Leu Arg Asp Gln
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                                      10
                                                           15
Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
             20
                                  25
                                                       30
Ala Arg Val Leu Ala Ile Glu Arg Tyr Leu Arg Asp Gln
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Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln 20 25 30

Ala Arg Val Leu 35

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Arg Tyr Leu Arg Asp Gln 35

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Gln Gln His Leu Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln 20 25 30

Ala Arg Val Leu Ala Val Glu Ser Tyr Leu Lys Asp Gln
35 40 45

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                                      10
                                                           15
Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Val Leu Ala Val Glu
                                  25
Ser Tyr Leu Lys Asp Gln
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Gln Gln His Leu Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
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Ala Arg Val Leu Ala Leu Glu Arg Tyr Leu Arg Asp Gln

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Arg Tyr Leu Arg Asp Gln
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  1
                                                           15
Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Val Lys Gln Leu Gln
             20
                                  25
Ala Arg Val Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln
         35
                              40
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1589.0160001Seq. Listing.txt
1 5 10 15
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Gln Gln His Met Leu Gln Leu Thr Val Trp Gly Val Lys Gln Leu Gln
20 25 30

Ala Arg Val Leu 35

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1 5 10 15

Thr Val Trp Gly Val Lys Gln Leu Gln Ala Arg Val Leu Ala Val Glu 20 25 30

Arg Tyr Leu Lys Asp Gln 35

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Gln Gln His Leu Leu Lys Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
20 25 30

Ala Arg Val Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln
35 40 45

<210> 33

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1589.0160001Seq. Listing.txt
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Gln Gln His Leu Leu Lys Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
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Ala Arg Val Leu
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Ser Asn Leu Leu Lys Ala Ile Glu Ala Gln Gln His Leu Leu Lys Leu
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                                                          15
Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Val Leu Ala Val Glu
             20
                                                      30
Arg Tyr Leu Lys Asp Gln
         35
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10

15

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Gln Gln His Leu Leu Gln Leu Ser Ile Trp Gly Ile Lys Gln Leu Gln
20 25 30

Ala Lys Val Leu Ala Ile Glu Arg Tyr Leu Arg Asp Gln
35 40 45

<210> 36

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Ser Gly Ile Val Gln Gln Gln Asn Ile Leu Leu Arg Ala Ile Glu Ala 1 5 10 15

Gln Gln His Leu Leu Gln Leu Ser Ile Trp Gly Ile Lys Gln Leu Gln
20 25 30

Ala Lys Val Leu 35

<210> 37

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1 5 10 15

Ser Ile Trp Gly Ile Lys Gln Leu Gln Ala Lys Val Leu Ala Ile Glu 20 25 30

Arg Tyr Leu Arg Asp Gln 35

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                                                          15
Gln Gln Leu Leu Arg Leu Ser Xaa Trp Gly Ile Arg Gln Leu Arg
             20
                                 25
Ala Arg Leu Leu Ala Leu Glu Thr Leu Leu Gln Asn Gln
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                             40
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Gln Gln Leu Leu Arg Leu Ser Xaa Trp Gly Ile Arg Gln Leu Arg
                                 25
Ala Arg Leu
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Ser Xaa Trp Gly Ile Arg Gln Leu Arg Ala Arg Leu Leu Ala Leu Glu 20 25 30

Thr Leu Leu Gln Asn Gln 35

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1 5 10 15

Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu 20 25 30

Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe 35 40 45

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1 5 10 15

Thr Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Asp 20 25 30

Leu Leu Ala Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe 35 40 45

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Thr Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Asp
                                  25
Leu Leu
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Tyr Thr Gly Leu Ile Tyr Thr Leu Ile Glu Glu Ser Gln Asn Gln Gln
  1
                  5
                                                          15
Glu Lys Asn Glu Gln Asp Leu Leu Ala Leu Asp Lys Trp Ala Ser Leu
             20
                                  25
                                                      30
Trp Asn Trp Phe
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Trp Met Glu Trp Glu Arg Glu Ile Asp Asn Tyr Thr Ser Glu Ile Tyr
Thr Leu Ile Glu Glu Ser Gln Asn Gln Glu Lys Asn Glu Gln Glu
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Page 19

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Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe 35 40 45

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Trp Met Glu Trp Glu Arg Glu Ile Asp Asn Tyr Thr Ser Glu Ile Tyr 1 5 10 15

Thr Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu 20 25 30

Leu Leu

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Glu Lys Asn Glu Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu 20 25 30

Trp Asn Trp Phe 35

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Asp Leu Leu Glu Lys Ser Gln Thr Gln Gln Glu Lys Asn Glu Lys Glu
20 25 30

Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe 35 40 45

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Asp Leu Leu Glu Lys Ser Gln Thr Gln Gln Glu Lys Asn Glu Lys Glu
20 25 30

Leu Leu

<210> 50

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Tyr Thr Asp Tyr Ile Tyr Asp Leu Leu Glu Lys Ser Gln Thr Gln Gln 1 5 10 15

Glu Lys Asn Glu Lys Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu 20 25 30

Trp Asn Trp Phe 35

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Trp Ile Gln Trp Asp Arg Glu Ile Ser Asn Tyr Thr Gly Ile Ile Tyr
1 5 10 15

Arg Leu Leu Glu Glu Ser Gln Asn Gln Gln Glu Asn Asn Glu Lys Asp 20 25 30

Leu Leu Ala Leu Asp Lys Trp Gln Asn Leu Trp Ser Trp Phe 35 40 45

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Trp Ile Gln Trp Asp Arg Glu Ile Ser Asn Tyr Thr Gly Ile Ile Tyr
1 5 10 15

Arg Leu Leu Glu Glu Ser Gln Asn Gln Gln Glu Asn Asn Glu Lys Asp 20 25 30

Leu Leu

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Leu Leu

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Glu Lys Asn Glu Lys Asp Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu 20 25 30

Trp Asn Trp Phe 35

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Trp Met Glu Trp Gln Lys Glu Ile Ser Asn Tyr Ser Asn Glu Val Tyr
1 5 10 15

Arg Leu Ile Glu Lys Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Gly
20 25 30

Leu Leu Ala Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe 35 40 45

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Arg Leu Ile Glu Lys Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Gly
             20
                                  25
Leu Leu
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                  5
                                      10
                                                          15
Glu Lys Asn Glu Gln Gly Leu Leu Ala Leu Asp Lys Trp Ala Ser Leu
             20
                                                      30
Trp Asn Trp Phe
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Trp Ile Gln Trp Asp Arq Glu Ile Ser Asn Tyr Thr Gln Gln Ile Tyr
                                      10
                                                          15
Ser Leu Ile Glu Glu Ser Gln Asn Gln Glu Lys Asn Glu Gln Asp
                                  25
Leu Leu Ala Leu Asp Asn Trp Ala Ser Leu Trp Thr Trp Phe
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Page 25

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<400> 61

Trp Ile Gln Trp Asp Arg Glu Ile Ser Asn Tyr Thr Gln Gln Ile Tyr 15

Ser Leu Ile Glu Glu Ser Gln Asn Gln Glu Lys Asn Glu Gln Asp 20 30 25

Leu Leu

<210> 62

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 62

Tyr Thr Gln Gln Ile Tyr Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln

Glu Lys Asn Glu Gln Asp Leu Leu Ala Leu Asp Asn Trp Ala Ser Leu 25 30

Trp Thr Trp Phe 35

<210> 63

<211> 46

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

Page 26

<400> 63

Trp Met Glu Trp Asp Arg Gln Ile Asp Asn Tyr Thr Glu Val Ile Tyr 1 5 10 15

Arg Leu Glu Leu Ser Gln Thr Gln Gln Gln Gln Asn Glu Gln Asp 20 25 30

Leu Leu Ala Leu Asp Lys Trp Asp Ser Leu Trp Asn Trp Phe 35 40 45

<210> 64

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 64

Trp Met Glu Trp Asp Arg Gln Ile Asp Asn Tyr Thr Glu Val Ile Tyr 1 5 10 15

Arg Leu Glu Leu Ser Gln Thr Gln Gln Gln Gln Asn Glu Gln Asp
20 25 30

Leu Leu

<210> 65

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 65

Tyr Thr Glu Val Ile Tyr Arg Leu Leu Glu Leu Ser Gln Thr Gln Gln
1 10 15

Glu Gln Asn Glu Gln Asp Leu Leu Ala Leu Asp Lys Trp Asp Ser Leu 20 25 30

Trp Asn Trp Phe

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<210> 66
<211> 47
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
<400> 66
Trp Ile Gln Trp Glu Arg Glu Ile Asn Asn Tyr Thr Gly Ile Ile Tyr
Ser Leu Ile Glu Glu Ala Gln Asn Gln Glu Asn Asn Glu Lys Asp
Leu Leu Ala Leu Asp Lys Trp Thr Asn Leu Trp Asn Trp Phe Asn
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<210> 67
<211> 34
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 67
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                                                          15
                                      10
Ser Leu Ile Glu Glu Ala Gln Asn Gln Glu Asn Asn Glu Lys Asp
             20
                                                      30
Leu Leu
<210> 68
<211> 37
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic

<220>

<400> 68

Tyr Thr Gly Ile Ile Tyr Ser Leu Ile Glu Glu Ala Gln Asn Gln Gln 1 5 10 15

Glu Asn Asn Glu Lys Asp Leu Leu Ala Leu Asp Lys Trp Thr Asn Leu 20 25 30

Trp Asn Trp Phe Asn 35

<210> 69

<211> 46

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 69

Trp Gln Gln Trp Asp Glu Lys Val Arg Asn Tyr Ser Gly Val Ile Phe 1 5 10 15

Gly Leu Ile Glu Gln Ala Gln Glu Gln Gln Asn Thr Asn Glu Lys Ser 20 25 30

Leu Leu Glu Leu Asp Gln Trp Asp Ser Leu Trp Ser Trp Phe 35 40 45

<210> 70

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 70

Trp Gln Gln Trp Asp Glu Lys Val Arg Asn Tyr Ser Gly Val Ile Phe 1 5 10 15

Gly Leu Ile Glu Gln Ala Gln Glu Gln Gln Asn Thr Asn Glu Lys Ser 20 25 30

Leu Leu

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<210> 71
<211> 36
<212> PRT
<213> Artificial Sequence
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Tyr Ser Gly Val Ile Phe Gly Leu Ile Glu Gln Ala Gln Glu Gln
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                                                          15
Asn Thr Asn Glu Lys Ser Leu Leu Glu Leu Asp Gln Trp Asp Ser Leu
             20
                                  25
Trp Ser Trp Phe
         35
<210> 72
<211> 46
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
<400> 72
Trp Gln Glu Trp Asp Arg Gln Ile Ser Asn Ile Ser Ser Thr Ile Tyr
Glu Glu Ile Gln Lys Ala Gln Val Gln Glu Gln Asn Glu Lys Lys
Leu Leu Glu Leu Asp Glu Trp Ala Ser Ile Trp Asn Trp Leu
         35
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<210> 73
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 73

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Trp Gln Glu Trp Asp Arg Gln Ile Ser Asn Ile Ser Ser Thr Ile Tyr
                                      10
Glu Glu Ile Gln Lys Ala Gln Val Gln Glu Gln Asn Glu Lys Lys
                                  25
Leu Leu
<210> 74
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
<400> 74
Ile Ser Ser Thr Ile Tyr Glu Glu Ile Gln Lys Ala Gln Val Gln Gln
Glu Gln Asn Glu Lys Lys Leu Leu Glu Leu Asp Glu Trp Ala Ser Ile
             20
                                  25
Trp Asn Trp Leu
         35
<210> 75
<211> 465
<212> DNA
<213> Artificial Sequence
<220>
<221> CDS
<222> (34)..(357)
<220>
<223> Description of Artificial Sequence: Synthetic
<400> 75
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                                     Met Gly Ile Val Gln Gln Gln
aac aat ttg ctg agg gct att gag gcg caa cag cac ctg ctg cag ctg
                                                                   102
                                Page 31
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Asn	Asn	Leu 10	Leu	Arg	Ala	Ile	Glu 15	Ala	Gln	Gln	His	Leu 20	Leu	Gln	Leu	
acc	gta	tgg	ggc	atc	aag	cag	ctg	cag	gca	cgc	atc	ctg	gct	gtt	gaa	150
Thr	Val 25	Trp	Gly	Ile	Lys	Gln 30	Leu	Gln	Ala	Arg	Ile 35	Leu	Ala	Val	Glu	
cgc	tac	ctg	aag	gat	caa	ggc	ggc	ggc	tca	ggc	gcc	ggc	tca	gag	tgg	198
Arg 40	Tyr	Leu	Lys	Asp	Gln 45	Gly	Gly	Gly	Ser	Gly 50	Ala	Gly	Ser	Glu	Trp 55	
gac	aga	gaa	att	aac	aat	tac	aca	agc	tta	ata	cac	tcc	tta	att	gaa	246
Asp	Arg	Glu	Ile	Asn 60	Asn	Tyr	Thr	Ser	Leu 65	Ile	His	Ser	Leu	Ile 70	Glu	
gaa	tcg	caa	aac	cag	caa	gaa	aag	aat	gaa	caa	gaa	tta	ttg	gaa	tta	294
Glu	Ser	Gln	Asn 75	Gln	Gln	Glu	Lys	Asn 80	Glu	Gln	Glu	Leu	Leu 85	Glu	Leu	
gat	aaa	tgg	gca	agt	ttg	tgg	aat	tgg	ttt	gaa	ttc	atc	gat	gat	atc	342
Asp	Lys	Trp 90	Ala	Ser	Leu	Trp	Asn 95	Trp	Phe	Glu	Phe	Ile 100	Asp	Asp	Ile	
aga	tcc	ggc	tgc	taa	caaa	agcco	cga a	aagga	aagct	g ag	gttt	ggct	gctg	gcca	cccg	397
Arg	Ser 105	Gly	Cys													
ctga	agcaa	ata a	actaç	gcata	aa co	ccctt	-gggg	g gcd	ctcta	aaac	gggt	cctt	gag g	gggti	ttttg	457
ctto	gaaaq	7														465

<210> 76

<211> 107

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence:Synthetic

<400> 76

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Met Gly Ile Val Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala
Gln Gln His Leu Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln
Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln Gly Gly
                             40
Ser Gly Ala Gly Ser Glu Trp Asp Arg Glu Ile Asn Asn Tyr Thr Ser
Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn
                                         75
Glu Gln Glu Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp
                 85
                                     90
                                                          95
Phe Glu Phe Ile Asp Asp Ile Arg Ser Gly Cys
            100
                                105
<210> 77
<211> 197
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
<400> 77
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tgetttetag aacegggtge geataaaaat geateaegee tatagegeta gageegetge 180
attaaatgaa tcggcca
                                                                   197
<210> 78
<211> 18
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
<400> 78
Tyr Thr Ser Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln
                                     10
                                                          15
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<210> 79
<211> 18
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
<400> 79
Glu Glu Ser Gln Asn Gln Glu Lys Asn Glu Gln Glu Leu Leu Glu
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                                      10
Leu Asp
<210> 80
<211> 13
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 80
Leu Glu Leu Asp Lys Trp Ala Ser Leu Trp Asn Trp Phe
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<210> 81
<211> 26
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
<400> 81
Gly Gly Gly Cys Cys Ala Thr Ala Thr Gly Gly Gly Thr Ala Thr
Thr Gly Thr Thr Cys Ala Gly Cys Ala Gly
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<210> 82
<211> 44
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
Gly Gly Cys Cys Gly Gly Cys Gly Cys Cys Thr Gly Ala Gly Cys
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Cys Gly Cys Cys Gly Cys Cys Thr Thr Gly Ala Thr Cys Cys Thr Thr
Cys Ala Gly Gly Thr Ala Gly Cys Gly Thr Thr Cys
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<210> 83
<211> 44
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 83
Gly Gly Cys Cys Gly Gly Cys Gly Cys Gly Gly Cys Thr Cys
Ala Gly Ala Gly Thr Gly Gly Gly Ala Cys Ala Gly Ala Gly Ala Ala
Ala Thr Thr Ala Ala Cys Ala Ala Thr Thr Ala Cys
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<210> 84
<211> 41
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 84

Gly Gly Cys Cys Gly Ala Ala Thr Thr Cys Thr Thr Ala Ala Ala 1 5 10 15

Ala Cys Cys Ala Ala Thr Thr Cys Cys Ala Cys Ala Ala Cys Thr 20 25 30

Thr Gly Cys Cys Cys Ala Thr Thr Thr 35 40